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(54) Title: METHOD OF DETERMINING THE FUNCTION OF NUCLEOTIDE SEQUENCES AND THE PROTEINS THEY ENCODE BY TRANSFECTING THE SAME INTO A HOST

### (57) Abstract

The present invention provides methods for rapidly determining the function of nucleic acid sequences by transfecting the same into a host organism to effect expression. Phenotypic and biochemical changes produced thereby are then analyzed to ascertain the function of the nucleic acids which have been transfected into the host organism. The invention also provides methods for silencing endogenous genes by transfecting hosts with nucleic acid sequences to effect expression of the same. The present invention also provides methods for selecting desired functions of RNAs and proteins by the use of virus vectors to express libraries of nucleic acid sequence variants. Moreover, the present invention provides methods for inhibiting an endogenous protease of a plant host.

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Inte. onal Application No PCT/US 99/01164

CLASSIFICATION OF SUBJECT MATTER C12N15/10 C12N15/67 C12N15/83 C12Q1/68 C12N15/82 //A01H3/00 According to International Patent Classification (IPC) or to both national classification and IPC **B. FIELDS SEARCHED** Minimum documentation searched (classification system followed by classification symbols) IPC 6 C12N Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practical, search terms used) C. DOCUMENTS CONSIDERED TO BE RELEVANT Category of Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No. X SABLOWSKI, R.W.M., ET AL.: "expression of 1-18, a flower-specific Myb protein in leaf 23-28, cells using a viral vector causes ectopic 56-58 activation of a target promoter" PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE USA. vol. 92, July 1995, pages 6901-6905, XP002118431 see the whole document X WO 95 34668 A (BIOSOURCE TECH INC) 21 1-18, December 1995 21-27, 29-33, 35-37, 56-58 see the whole document Further documents are listed in the continuation of box C. Patent family members are listed in annex. Special categories of cited documents: "T" later document published after the international filing date \*A\* document defining the general state of the art which is not considered to be of particular relevance or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention \*E\* earlier document but published on or after the international \*X\* document of particular relevance; the claimed invention filing date cannot be considered novel or cannot be considered to "L" document which may throw doubts on priority claim(s) or involve an inventive step when the document is taken alone which is cited to establish the publication date of another citation or other special reason (as specified) "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such docu-\*O\* document referring to an oral disclosure, use, exhibition or other means ments, such combination being obvious to a person skilled document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of the actual completion of the international search Date of mailing of the international search report 2 7. 01. 2000 15 October 1999 Name and mailing address of the ISA **Authorized officer** European Patent Office, P.B. 5818 Patentiaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni, Holtorf, S Fax: (+31-70) 340-3016

Intel anal Application No
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(	WO 91 01375 A (ICI PLC) 7 February 1991	51-53, 56-58		
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PCT/US 99/01164

Box I	Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This Inte	rnational Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1.	Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
2.	Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
з. 🗌	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II	Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This Inte	ernational Searching Authority found multiple inventions in this international application, as follows:
	see additional sheet
1.	As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2.	As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3.	As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. X	No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:  1, 51-53, 56-58 (partially); 2-59, 117 (completely)
Remar	k on Protest  The additional search fees were accompanied by the applicant's protest.  No protest accompanied the payment of additional search fees.

# FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

1. Claims: 1,51-53, 56-58 partially, 2-50,117 completely

A method to determine the function of a nucleic acid sequence or members of a cDNA- or EST-library in an organism by utilizing a viral expression vector and observing changes which result from the expression of the nucleic acid; a method of silencing endogenous genes using such viral vector; a method for the construction of a cDNA library.

2. Claims: 51-53, 56-58 partially, 54,55,59-74 completely

A method to determine the function of a nucleic acid sequence by altering the genome of a host organism by transposon mutagenesis and observing changes which result from the subsequent expression of the nucleic acid of interest.

3. Claims: 75-77 completely

A method to identify a gene product target of an inhibitor by identifying the gene product that corresponds to a conditional lethal mutation.

4. Claims: 78-92 completely

A method for constructing an infectious viral vector and method for infecting a plant host with such a vector.

5. Claims: 93-97 completely

A method for inhibiting an endogenous protease of a plant by applying a compound that induces the production of an inhibitor of said protease.

6. Claim: 98 completely

A method for improving the expression of an exogenous nucleic acid sequence in a plant comprising interspecific hybridisation.

7. Claims: 99-114 completely

A method for optimizing the function of a nucleic acid sequence in a plant by expressing variant forms of the nucleic acid sequence utilizing a viral expression vector.

8. Claim: 115 completely

# FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

A method of increasing the representation of a nucleic acid sequence in a viral expression library by propagating said library in the absence of E. coli.

9. Claim: 1 partially; 116 completely

A method to determine the function of a gene wherein one or more reporter genes are fused to one or more promoters in a viral expression vector.

information on patent family members

Inte ional Application No
PCT/US 99/01164

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